sampleMap

	A	B	C
1	assay	primary	colname
2	RNASeq2GeneNorm	A5J1	A5J2-01A
3	RNASeq2GeneNorm	A5J2	A5J3-01A
4	RNASeq2GeneNorm	A5J3	A5J5-01A
5	Mutations	A5J1	A5J2-11D
6	Mutations	A5J2	A5J3-11D
7	Mutations	A5J3	A5J4-11D

patient IDs as defined in the **colData**

patientID	age	tumor_tissu	pathologic_stage	gender
A5J1	58	adrenal	stage ii	male
A5J2	44	adrenal	stage iv	female
A5J3	23	adrenal	stage iii	female

gene_symbo	A5J2-01A	A5J3-01A	A5J5-01A
DIRAS3	1487.0317	9.6631	18.9602
MAPK14	778.5783	2823.6469	1061.7686
YAP1	1009.6061	2305.059	1561.2502
CDKN1B	2101.3449	4248.9584	1348.541
ERBB2	651.2968	246.4098	90.0607
G6PD	1456.7723	1715.6081	3571.0265
KDR	349.6638	1246.141	486.4465

column names of **RNAseq** results

column names of **mutation** assay results

gene_symb	A5J2-11D	A5J3-11D	A5J4-11D
DIRAS3	0	0	0
G6PD	0	0	0
KDR	0	0	0
ERBB3	0	0	0
AKT1S1	0	0	0
RPS6KA1	0	0	0
MRE11A	0	0	0

the results of the different assays; will be stored in `experiment`